GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: May 30, 2003, 11:31:08 ; Search time 159 Seconds
(without alignments)
7641.845 Million cell updates/sec

Title: US-08-153-397A-1
Perfect score: 3962
Sequence: 1 CGGGCCTGAGACTGGGGTGA.....AAAAAAAAAAACCGGAATTC 3962

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 441362 seqs, 153338381 residues

Total number of hits satisfying chosen parameters: 882724

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
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3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
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5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	3962	100.0	3962	1	US-08-336-343A-1	Sequence 1, Appli
2	3451	87.1	3637	1	US-08-445-640-3	Sequence 3, Appli
3	3451	87.1	3637	3	US-08-170-558-3	Sequence 3, Appli
4	3451	87.1	3637	3	US-08-447-314-3	Sequence 3, Appli
5	3451	87.1	3637	3	US-08-445-461-3	Sequence 3, Appli
6	1192.2	30.1	1197	1	US-08-445-640-7	Sequence 7, Appli
7	1192.2	30.1	1197	3	US-08-170-558-7	Sequence 7, Appli
8	1192.2	30.1	1197	3	US-08-447-314-7	Sequence 7, Appli
9	1192.2	30.1	1197	3	US-08-445-461-7	Sequence 7, Appli
10	642	16.2	3157	1	US-08-336-343A-3	Sequence 3, Appli
c 11	642	16.2	3157	1	US-08-336-343A-5	Sequence 5, Appli
12	639.8	16.1	3120	1	US-08-456-647B-19	Sequence 19, Appl
13	639.8	16.1	3120	2	US-08-237-401A-19	Sequence 19, Appl
14	182.2	4.6	2820	1	US-08-286-305A-4	Sequence 4, Appli
15	182.2	4.6	2820	2	US-08-441-104A-4	Sequence 4, Appli
16	182.2	4.6	2820	2	US-08-440-816A-4	Sequence 4, Appli
17	182.2	4.6	2820	4	US-09-417-381A-4	Sequence 4, Appli
18	180.6	4.6	2301	1	US-08-306-691B-23	Sequence 23, Appl
19	180.6	4.6	2301	5	PCT-US93-06251-78	Sequence 78, Appl
20	180.6	4.6	3060	1	US-08-286-305A-6	Sequence 6, Appli
21	180.6	4.6	3060	2	US-08-441-104A-6	Sequence 6, Appli
22	180.6	4.6	3060	2	US-08-440-816A-6	Sequence 6, Appli
23	180.6	4.6	3060	4	US-09-417-381A-6	Sequence 6, Appli
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25	180.6	4.6	3194	2	US-08-286-846A-1	Sequence 1, Appli
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27	180.6	4.6	3194	3	US-08-444-622A-1	Sequence 1, Appli
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29	180.6	4.6	3194	3	US-09-156-923-1	Sequence 1, Appli
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33	178.6	4.5	2940	1	US-08-286-305A-8	Sequence 8, Appli
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36	178.6	4.5	2940	4	US-09-417-381A-8	Sequence 8, Appli
37	163	4.1	2463	1	US-08-339-578-1	Sequence 1, Appli
38	158	4.0	4092	2	US-08-469-537A-106	Sequence 106, App
39	147.2	3.7	3398	5	PCT-US95-08493-12	Sequence 12, Appl
40	139	3.5	2208	5	PCT-US95-08493-1	Sequence 1, Appli
41	139	3.5	2580	5	PCT-US95-08493-18	Sequence 18, Appl
42	139	3.5	2604	5	PCT-US95-08493-20	Sequence 20, Appl
43	138.4	3.5	2376	1	US-07-912-952-3	Sequence 3, Appli
44	135.2	3.4	4149	2	US-08-737-715-1	Sequence 1, Appli
45	130.4	3.3	2869	1	US-08-374-834-2	Sequence 2, Appli

ALIGNMENTS

RESULT 1
US-08-336-343A-1
; Sequence 1, Application US/08336343A
; Patent No. 5677144
; GENERAL INFORMATION:
; APPLICANT: Ullrich, Axel
; APPLICANT: Alves, Frauke
; TITLE OF INVENTION: CCK-2, A No. 5677144el Receptor Tyrosine Kinase
; NUMBER OF SEQUENCES: 43
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/336,343A
; FILING DATE: 08-NOV-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Coruzzi, Laura A.
; REGISTRATION NUMBER: 30,742
; REFERENCE/DOCKET NUMBER: 7683-065
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3962 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: unknown
; MOLECULE TYPE: cDNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 321..3077
US-08-336-343A-1

Query Match 100.0%; Score 3962; DB 1; Length 3962;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 3962; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db      61 CTGCTCTCGGGAGCCGCCTCCCGACACCCGAGCCCCGCCGGCGCCTCCCGCTCCCGGCTC 120

Qy     121 CCGGCTCCTGGCTCCCTCCGCCTCCCCCGCCCCTCGCCCCGCCGCCGAAGAGGCCCCGCT 180
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Qy     181 CCCGGGTCGGACGCCTGGGTCTGCCGGGAAGAGCGATGAGAGGTGTCTGAAGGTGGCTAT 240
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Db       2521 GGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAG 2580
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Qy       2641 CCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGA 2700
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Qy       2701 GCCGGAACCTCTATGCTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCC 2760
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Qy       2761 GCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGG 2820
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Qy       3301 TAGAAGCCCCTGTCGCCCACCCAGCTGGTCCTGTGGATGGGATCCTCTCCACCCTCCTCT 3360
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Db       3361 AGCCATCCCTTGGGGAAGGGTGGGGAGAAATATAGGATAGACACTGGACATGGCCCATTG 3420
Qy       3421 GAGCACCTGGGCCCCACTGGACAACACTGATTCCTGGAGAGGTGGCTGCGCCCCAGCTTC 3480
Db       3421 GAGCACCTGGGCCCCACTGGACAACACTGATTCCTGGAGAGGTGGCTGCGCCCCAGCTTC 3480
Qy       3481 TCTCTCCCTGTCACACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGA 3540
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Db       3601 CTCCTCCATCACCTGAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCC 3660
Qy       3661 CCACTTCCCACTTGCAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAG 3720
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Qy       3841 GGAGAGACACAGATTTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCT 3900
Db       3841 GGAGAGACACAGATTTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCT 3900
Qy       3901 GCACTAGGCAGGTAATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAAACCGGAAT 3960
Db       3901 GCACTAGGCAGGTAATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAAACCGGAAT 3960
Qy       3961 TC 3962
Db       3961 TC 3962
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RESULT 2
US-08-445-640-3
; Sequence 3, Application US/08445640
; Patent No. 5709858
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
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;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  patin (Genentech)
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/445,640
;       FILING DATE:  22-MAY-1995
;       CLASSIFICATION:  435
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/170558
;       FILING DATE:  20-DEC-1993
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/157563
;       FILING DATE:  23-NOV-1993
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Hasak, Janet E.
;       REGISTRATION NUMBER:  28,616
;       REFERENCE/DOCKET NUMBER:  8344C2
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  415/225-1896
;       TELEFAX:  415/952-9881
;       TELEX:  910/371-7168
;   INFORMATION FOR SEQ ID NO:  3:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  3637 bases
;       TYPE:  nucleic acid
;       STRANDEDNESS:  single
;       TOPOLOGY:  linear
US-08-445-640-3

  Query Match             87.1%;  Score 3451;  DB 1;  Length 3637;
  Best Local Similarity   97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

Qy        256 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 315
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Qy        436 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 495
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        197 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 256

Qy        496 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 555
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        257 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 316

Qy        556 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCTC 615
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db        317 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCTC 376

Qy        616 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 675
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        377 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 436

Qy        676 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 735
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        437 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 496

Qy        736 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 795
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Db        497 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 556

Qy        796 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTGC 855
              ||||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db        557 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTGC 616

Qy        856 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 915
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        617 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 676

Qy        916 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 975
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Db        677 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 736

Qy        976 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 1035
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Db        737 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 796

Qy       1036 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 1095
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        797 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 856

Qy       1096 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 1155
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        857 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 916

Qy       1156 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 1215
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        917 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 976

Qy       1216 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1275
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Db        977 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1036

Qy       1276 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1335
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Qy       1336 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1395
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Qy       1396 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1455
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Db        1157 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1216

Qy       1456 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1515
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Db        1217 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1276

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Db        1277 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCCATCC 1336

Qy       1576 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1635
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              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1397 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1456

Qy       1696 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1755
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Qy       1756 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1815
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1517 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1576

Qy       1816 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1875
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy       1876 CTTACGCCCGTCCCCCTCGAGGCCCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCA 1935
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Qy      1996 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 2055
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy      2056 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 2115
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1817 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 1876

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Qy      2236 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2295
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy      2296 CAGATGCCACCAAGAATGCCAGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGG 2355
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Qy      2356 TGAAGATCATGTCGAGGCTCAAGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGC 2415
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Db      2159 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2218

Qy      2476 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2535
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Db      2219 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2278

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Db      2339 GCATGCGCTATCTGGCCACACTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCC 2398

Qy      2656 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2715
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Db      2399 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2458

Qy      2716 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2775
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      2459 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2518

Qy      2776 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2835
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Db      2519 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2578

Qy      2836 TGTGGGAGGTGCTGATGCTCTGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGG 2895
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Db      2699 CGCCTGCCTGCCCGCAGGGCCTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTG 2758

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Db      2759 AGCAGCGACCACCCTTTTCCCAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGG 2818

Qy      3076 TGTGAATCACACATCCAGCTGCCCCTCCCTCAGGGAGTGATCCAGGGGAAGCCAGTGACA 3135
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Db      2819 TGTGAATCACACATCCAGCTGCCCCTCCCTCAGGGAGTGATCCAGGGGAAGCCAGTGACA 2878

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Qy      3436 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCG-CCCCAGCTTCTCTCTCCCTGTCAC 3494
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Qy      3495 ACACTGGACCCCACTGGCTGAGAATCTGGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3554
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Db      3153 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3212

Qy      3555 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3614
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Db      3213 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3272

Qy      3615 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACTTG 3674
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Db      3273 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACCTG 3332

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Qy      3735 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3794
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Db      3393 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3452

Qy      3795 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3854
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Qy      3855 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3914
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RESULT 3
US-08-170-558-3
; Sequence 3, Application US/08170558
; Patent No. 6001621
; GENERAL INFORMATION:
; APPLICANT: Godowski, Paul J.
; APPLICANT: Mark, Melanie R.
; APPLICANT: Scadden, David T.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Baron, Will F.
; TITLE OF INVENTION: Protein Tyrosine Kinases

```
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/170,558
;      FILING DATE:  20-DEC-1993
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3637 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-170-558-3

  Query Match             87.1%;  Score 3451;  DB 3;  Length 3637;
  Best Local Similarity   97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

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              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        377 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 436

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              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        437 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 496

Qy        736 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 795
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              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy        916 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 975
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Db        677 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 736

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Db        737 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 796

Qy       1036 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 1095
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Qy       1216 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1275
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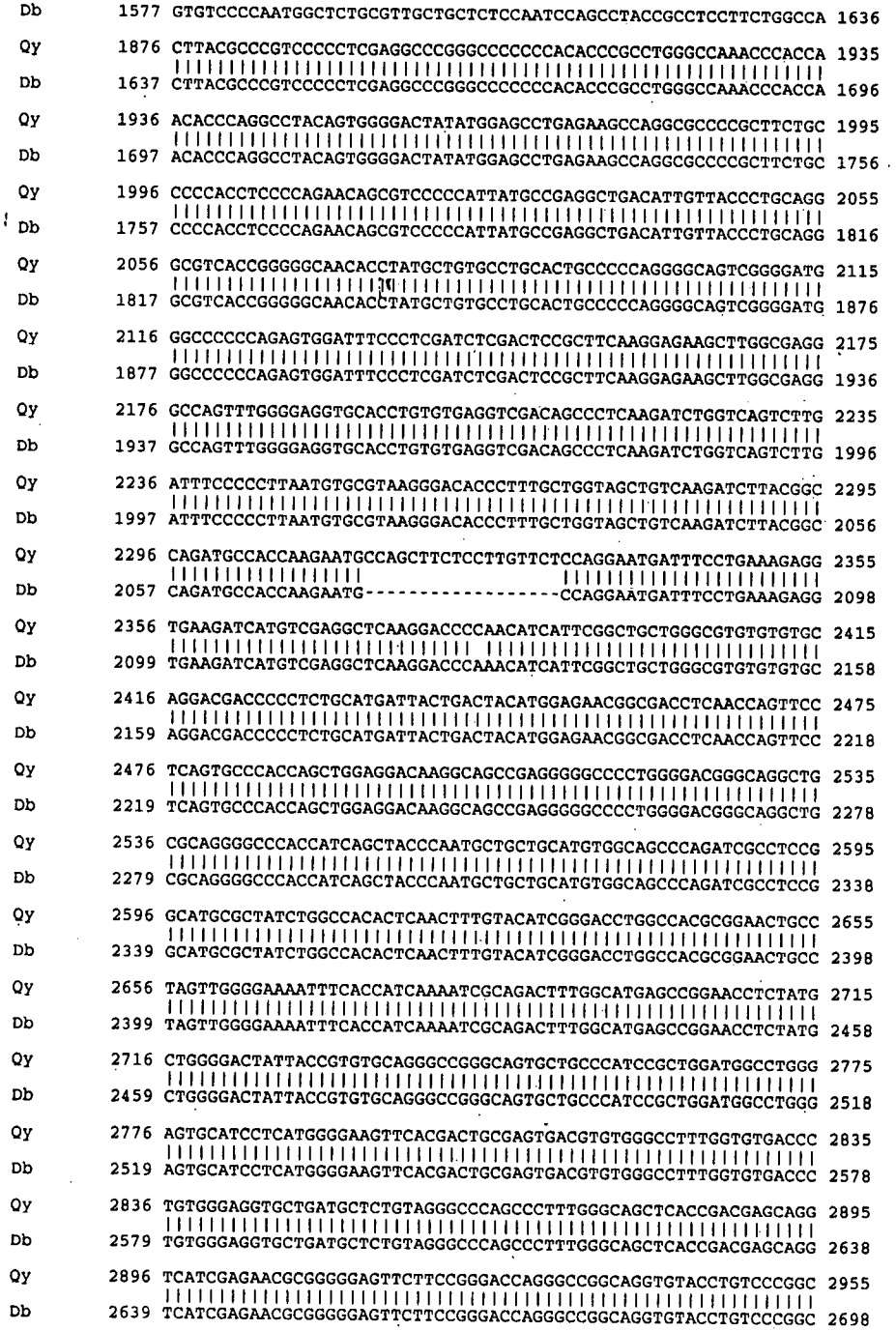

RESULT 4
US-08-447-314-3
; Sequence 3, Application US/08447314

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; Patent No. 6087144
;  GENERAL INFORMATION:
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/447,314
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1D2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3637 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-447-314-3

  Query Match            87.1%;  Score 3451;  DB 3;  Length 3637;
  Best Local Similarity  97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

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Db        137 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 196

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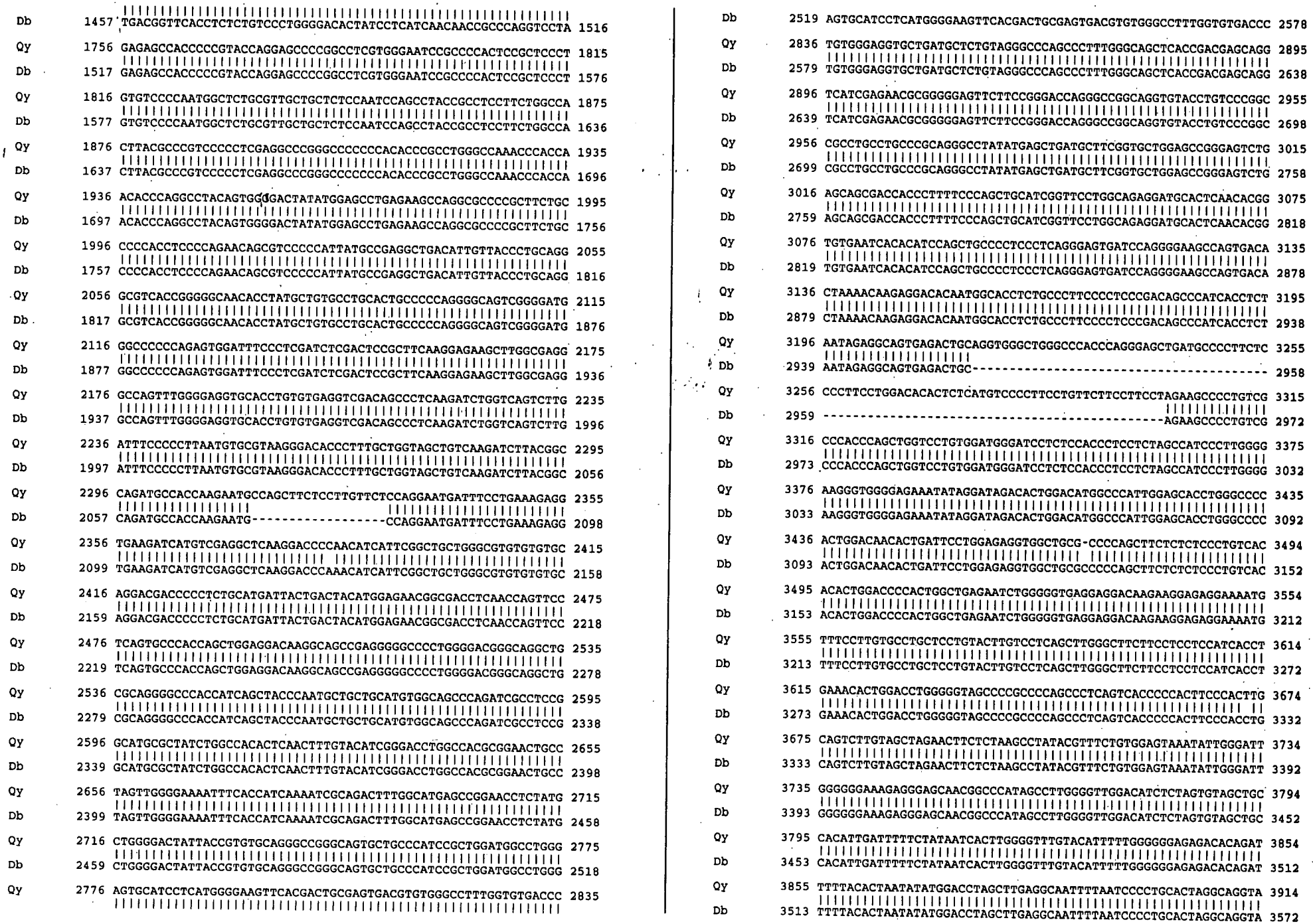
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RESULT 5
US-08-445-461-3
; Sequence 3, Application US/08445461
; Patent No. 6096527
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/445,461
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  530
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C3
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3637 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-445-461-3

  Query Match             87.1%;  Score 3451;  DB 3;  Length 3637;
  Best Local Similarity   97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

Qy        256 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 315
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         17 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 76

Qy        316 GAGCTATGGGACCAGAGGCCCTGTCATCTTTACTGCTGCTGCTCTTGGTGGCAAGTGGAG 375
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         77 GAGCTATGGGACCAGAGGCCCTGTCATCTTTACTGCTGCTGCTCTTGGTGGCAAGTGGAG 136

Qy        376 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 435
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        137 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 196

Qy        436 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 495
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        197 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 256

Qy        496 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 555
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        257 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 316

Qy        556 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCTC 615
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db        317 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCTC 376

Qy        616 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 675
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        377 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 436

Qy        676 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 735
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        437 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 496

Qy        736 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 795
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        497 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 556

Qy        796 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTGC 855
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||| ||
Db        557 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTGC 616

Qy        856 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 915
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        617 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 676

Qy        916 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 975
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        677 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 736

Qy        976 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 1035
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        737 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 796

Qy       1036 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 1095
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        797 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 856

Qy       1096 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 1155
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        857 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 916

Qy       1156 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 1215
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        917 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 976

Qy       1216 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1275
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        977 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1036

Qy       1276 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1335
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1037 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1096

Qy       1336 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1395
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1097 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1156

Qy       1396 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1455
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1157 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1216

Qy       1456 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1515
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        1217 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1276

Qy       1516 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCC 1575
              ||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
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Db       1277 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCCATCC 1336
Qy       1576 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1635
Db       1337 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1396
Qy       1636 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1695
Db       1397 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1456
Qy       1696 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1755
Db       1457 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1516
Qy       1756 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1815
Db       1517 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1576
Qy       1816 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1875
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Qy       1936 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1995
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Qy       1996 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 2055
Db       1757 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 1816
Qy       2056 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 2115
Db       1817 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 1876
Qy       2116 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 2175
Db       1877 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 1936
Qy       2176 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 2235
Db       1937 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 1996
Qy       2236 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2295
Db       1997 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2056
Qy       2296 CAGATGCCACCAAGAATGCCAGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGG 2355
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Qy       2416 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2475
Db       2159 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2218
Qy       2476 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2535
Db       2219 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2278
Qy       2536 CGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCG 2595
Db       2279 CGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCG 2338
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Qy       2656 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2715
Db       2399 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2458
Qy       2716 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2775
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Qy       2776 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2835
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Qy       2896 TCATCGAGAACGCGGGGGAGTTCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGC 2955
Db       2639 TCATCGAGAACGCGGGGGAGTTCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGC 2698
Qy       2956 CGCCTGCCTGCCCGCAGGGCCTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTG 3015
Db       2699 CGCCTGCCTGCCCGCAGGGCCTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTG 2758
Qy       3016 AGCAGCGACCACCCTTTTCCCAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGG 3075
Db       2759 AGCAGCGACCACCCTTTTCCCAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGG 2818
Qy       3076 TGTGAATCACACATCCAGCTGCCCCTCCCTCAGGGAGTGATCCAGGGGAAGCCAGTGACA 3135
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Qy       3136 CTAAAACAAGAGGACACAATGGCACCTCTGCCCTTCCCCTCCCGACAGCCCATCACCTCT 3195
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Qy       3196 AATAGAGGCAGTGAGACTGCAGGTGGGCTGGGCCCACCCAGGGAGCTGATGCCCCTTCTC 3255
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Qy       3256 CCCTTCCTGGACACACTCTCATGTCCCCTTCCTGTTCTTCCTTCCTAGAAGCCCCTGTCG 3315
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Qy       3316 CCCACCCAGCTGGTCCTGTGGATGGGATCCTCTCCACCCTCCTCTAGCCATCCCTTGGGG 3375
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Qy       3376 AAGGGTGGGGAGAAATATAGGATAGACACTGGACATGGCCCATTGGAGCACCTGGGCCCC 3435
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Qy       3436 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCG-CCCCAGCTTCTCTCTCCCTGTCAC 3494
Db       3093 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCGCCCCCAGCTTCTCTCTCCCTGTCAC 3152
Qy       3495 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3554
Db       3153 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3212
Qy       3555 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3614
Db       3213 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3272
Qy       3615 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACTTG 3674
Db       3273 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACCTG 3332
Qy       3675 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3734
Db       3333 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3392
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Qy       3735 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3794
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Db       3393 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3452

Qy       3795 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3854
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3453 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3512

Qy       3855 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3914
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3513 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3572

Qy       3915 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3953
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Db       3573 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3611
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RESULT 6
US-08-445-640-7
; Sequence 7, Application US/08445640
; Patent No. 5709858
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/445,640
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-445-640-7

  Query Match             30.1%;  Score 1192.2;  DB 1;  Length 1197;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-271;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy        375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
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Db          1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

Qy        435 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 494
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 120

Qy        495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy        555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db        181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy        615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

Qy        675 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 734
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

Qy        735 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 794
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        361 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 420

Qy        795 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTG 854
              ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||
Db        421 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTG 480

Qy        855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
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Db        481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

Qy        915 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 974
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Db        541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

Qy        975 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 1034
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Db        601 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 660

Qy       1035 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 1094
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        661 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 720

Qy       1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

Qy       1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

Qy       1215 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 1274
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        841 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 900

Qy       1275 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 1334
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        901 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 960

Qy       1335 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1394
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        961 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1020

Qy       1395 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1454
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Db       1021 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1080

Qy       1455 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1514
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1081 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1140
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Qy       1515 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCC 1571
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Db       1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197


RESULT 7
US-08-170-558-7
; Sequence 7, Application US/08170558
; Patent No. 6001621
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/170,558
;      FILING DATE:  20-DEC-1993
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-170-558-7

  Query Match             30.1%;  Score 1192.2;  DB 3;  Length 1197;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-271;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy        375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
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Db          1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

Qy        435 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 494
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Qy        495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
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Db        121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy        555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
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Db        181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy        615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
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Db        241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

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Db        301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

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Db        361 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 420

Qy        795 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTG 854
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Qy        855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
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Db        481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

Qy        915 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 974
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Db        541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

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Db        601 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 660

Qy       1035 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 1094
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Db        721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

Qy       1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
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Db        781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

Qy       1215 GGCGGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 1274
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Qy       1275 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 1334
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        901 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 960

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Db        961 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1020

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Db       1021 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1080

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Db       1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197


RESULT 8
US-08-447-314-7
; Sequence 7, Application US/08447314
; Patent No. 6087144
;  GENERAL INFORMATION:
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
```

```
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/447,314
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1D2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-447-314-7

  Query Match           30.1%;  Score 1192.2;  DB 3;  Length 1197;
  Best Local Similarity  99.7%;  Pred. No. 1.1e-271;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy        375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
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Db          1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

Qy        435 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 494
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Db         61 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 120

Qy        495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
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Db        121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy        555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
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Db        181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy        615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
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Db        241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

Qy        675 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 734
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

Qy        735 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 794
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Qy        855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

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Qy       1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
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Qy       1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
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Db        781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

Qy       1215 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 1274
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Db        841 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 900

Qy       1275 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 1334
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Db        901 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 960

Qy       1335 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1394
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Db       1081 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1140

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Db       1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197

RESULT 9
US-08-445-461-7
; Sequence 7, Application US/08445461
; Patent No. 6096527
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
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;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  patin (Genentech)
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/445,461
;       FILING DATE:  22-MAY-1995
;       CLASSIFICATION:  530
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/170558
;       FILING DATE:  20-DEC-1993
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/157563
;       FILING DATE:  23-NOV-1993
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Hasak, Janet E.
;       REGISTRATION NUMBER:  28,616
;       REFERENCE/DOCKET NUMBER:  854C3
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  415/225-1896
;       TELEFAX:  415/952-9881
;       TELEX:  910/371-7168
;   INFORMATION FOR SEQ ID NO:  7:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  1197 bases
;       TYPE:  nucleic acid
;       STRANDEDNESS:  single
;       TOPOLOGY:  linear
US-08-445-461-7

  Query Match             30.1%;  Score 1192.2;  DB 3;  Length 1197;
  Best Local Similarity   99.7%;  Pred. No. 1.1e-271;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy        375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
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Db          1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

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Db        241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

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Db        421 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTG 480

Qy        855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
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Qy        915 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 974
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Db        541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

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Qy       1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
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Db        721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

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Db        781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

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Qy       1395 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1454
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Qy       1515 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCC 1571
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Db       1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197


RESULT 10
US-08-336-343A-3
; Sequence 3, Application US/08336343A
; Patent No. 5677144
;  GENERAL INFORMATION:
;    APPLICANT:  Ullrich, Axel
;    APPLICANT:  Alves, Frauke
;    TITLE OF INVENTION:  CCK-2, A No. 5677144el Receptor Tyrosine Kinase
;    NUMBER OF SEQUENCES:  43
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Pennie & Edmonds
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/336,343A
;      FILING DATE:  08-NOV-1994
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Coruzzi, Laura A.
;      REGISTRATION NUMBER:  30,742
;      REFERENCE/DOCKET NUMBER:  7683-065
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 790-9090
;      TELEFAX:  (212) 869-9741/8864
```

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;      TELEX:  66141 PENNIE
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3157 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  both
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  370..2934
US-08-336-343A-3
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Query Match 16.2%; Score 642; DB 1; Length 3157;
Best Local Similarity 56.5%; Pred. No. 7.2e-142;
Matches 1534; Conservative 0; Mismatches 975; Indels 207; Gaps 10;

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Qy       349 TGCTGCTGCTCTTGGTGGCAAGTGGAGATGCTGACATGAAGGGACATTTTGATCCTGCCA 408
Db       395 TGGTGCTGTTCCTGCTGCTGCCTATCTTGAGTTCTGCAAAAGCTCAGGTTAATCCAGCTA 454

Qy       409 AGTGCCGCTATGCCCTGGGCATGCAGGACCGGACCATCCCAGACAGTGACATCTCTGCTT 468
Db       455 TATGCCGCTATCCTCTGGGCATGTCAGGAGGCCAGATTCCAGATGAGGACATCACAGCTT 514

Qy       469 CCAGCTCCTGGTCAGATTCCACTGCCGCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGG 528
Db       515 CCAGTCAGTGGTCAGAGTCCACAGCTGCCAAATATGGAAGGCTGGACTCAGAAGAAGGGG 574

Qy       529 ATGGGGCCTGGTGCCCCGCAGGGTCGGTGTTTCCCAAGGA---GGAGGAGTACTTGCAGG 585
Db       575 ATGGAGCCTGGTGCCCTGAGATTCCAGTGGAACCTGATGACCTGAAGGAGTTTCTGCAGA 634

Qy       586 TGGATCTACAACGACTCCACCTGGTGGCTCTGGTGGGCACCCAGGGACGGCATGCCGGGG 645
Db       635 TTGACTTGCACACCCTCCATTTTATCACTCTGGTGGGGACCCAGGGGCGCCATGCAGGAG 694

Qy       646 GCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGCT 705
Db       695 GTCATGGCATCGAGTTTGCCCCCATGTACAAGATCAATTACAGTCGGGATGGCACTCGCT 754

Qy       706 GGATGGGCTGGAAGGACCGCTGGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCTGAGG 765
Db       755 GGATCTCTTGGCGGAACCGTCATGGGAAACAGGTGCTGGATGGAAATAGTAACCCCTATG 814

Qy       766 GAGTGGTGCTGAAGGACCTTGGGCCCCCCATGGTTGCCCGACTGGTTCGCTTCTACCCCC 825
Db       815 ACATTTTCCTAAAGGACTTGGAGCCGCCCATTGTAGCCAGATTTGTCCGGTTCATTCCAG 874

Qy       826 GGGCTGACCGGGTCATGAGTGTCTGTCTGCGGGTAGAGCTCTATGGCTGCCTCTGGAGGG 885
Db       875 TCACCGACCACTCCATGAATGTGTGTATGAGAGTGGAGCTTTACGGCTGTGTCTGGCTAG 934

Qy       886 ATGGACTCCTGTCTTACACCGCCCCTGTGGGGCAGACAATGTATTTATCTGAGG------ 939
Db       935 ATGGCTTGGTGTCTTACAATGCTCCAGCTGGGCAGCAGTTTGTACTCCCTGGAGGTTCCA 994

Qy       940 CCGTGTACCTCAACGACTCCACCTATGACGGACATACCGTGGGCGGACTGCAGTATGGGG 999
Db       995 TCATTTATCTGAATGATTCTGTCTATGATGGAGCTGTTGGATACAGCATGACAGAAGGG- 1053

Qy      1000 GTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGGATGACTTTAGGAAGAGTCAGGAGC 1059
Db      1054 --CTAGGCCAATTGACCGATGGTGTGTCTGGCCTGGACGATTTCACCCAGACCCATGAAT 1111

Qy      1060 TGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGAGCAACCACAGCTTCTCCAGTGGCT 1119
Db      1112 ACCACGTGTGGCCCGGCTATGACTATGTGGGCTGGCGGAACGAGAGTGCCACCAATGGCT 1171

Qy      1120 ATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCACT 1179
Db      1172 ACATTGAGATCATGTTTGAATTTGACCGCATCAGGAATTTCACTACCATGAAGGTCCACT 1231

Qy      1180 GTAACAACATGCACACGCTGGGAGCCCGTCTGCCTGGCGGGGTGGAATGTCGCTTCCGGC 1239
Db      1232 GCAACAACATGTTTGCTAAAGGTGTGAAGATCTTTAAGGAGGTACAGTGCTACTTCCGCT 1291

Qy      1240 GTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTGG 1299
Db      1292 CTG---AAGCCAGTGAGTGGGAACCTAATGCCATTTCCTTCCCCCTTGTCCTGGATGACG 1348

Qy      1300 GGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTGC 1359
Db      1349 TCAACCCCAGTGCTCGGTTTGTCACGGTGCCTCTCCACCACCGAATGGCCAGTGCCATCA 1408

Qy      1360 AGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCTTCAGCGAAATCTCCTTCATCTCTG 1419
Db      1409 AGTGTCAATACCATTTTGCAGATACCTGGATGATGTTCAGTGAGATCACCTTCCAATCAG 1468

Qy      1420 ATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCACCTTCCCGCCAGCCCCCTGGTGGC 1479
Db      1469 ATGCTGCAATGTACAACAACTCTGAAGCCCTGCCCACCTCTCC---------------- 1511

Qy      1480 CGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGGAGCTGGAGCCCAGAGGCCAGCAGC 1539
Db      1512 ----------------------------------TATGGCACCCACAACCTATGATC 1534

Qy      1540 CCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCCTCATCGGCTGCCTGGTGGCCATCA 1599
Db      1535 CAATGCTTAAAGTTGATGACAGCAACACTCGGATCCTGATTGGCTGCTTGGTGGCCATCA 1594

Qy      1600 TCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCTGGCGGCTGCACTGGCGCAGGCTCC 1659
Db      1595 TCTTTATCCTCCTGGCCATCATTGTCATCATCCTCTGGAGGCAGTTCTGGCAGAAAATGC 1654

Qy      1660 TCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGCTGACGGTTCACCTCTCTGTCCCTG 1719
Db      1655 TGGAGAAGGCTTCTCGGAGGATGCTGGATGATGAAATGACAGTCAGCCTTTCCCTGCCAA 1714

Qy      1720 GGGACACTATCCTCATCAACAACCGCCCAGGTCCTAGAGAGCCACCCCCGTACCAGGAGC 1779
Db      1715 GTGATTCTAGCATGTTCAACAAT----------------------------AACCGCTCCT 1747

Qy      1780 CCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCTGTGTCCCCAATGGCTCTGCGTTGC 1839
Db      1748 CATCACCTAGTGAACAAGGGTCCAACTCGACTTACGATCGCATCTTT------------ 1794

Qy      1840 TGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCACTTACGCCCGTCCCCCTCGAGGCC 1899
Db      1795 ----------------------------------------------------CCCCTTCGCC 1804

Qy      1900 CGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCAACACCCAGGCCTACAGTGGGGACT 1959
Db      1805 CTGACTACCAGGAGCCATCCAGGCTGATACGAAAACTCCCAGAATTTGCTCCAGGGGAGG 1864

Qy      1960 ATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGCCCCCACCTCCCCAGAACAGCGTCC 2019
Db      1865 AGGAGTCAGGCTGCAGCGGTGTTGTGAAGCCAGTCCAGCCCAGTGGCCCTGAGGGGGTGC 1924

Qy      2020 CCCATTATGCCGAGGCTGACATTGTTACCCTGCAGGGCGTCACCGGGGGCAACACCTATG 2079
Db      1925 CCCACTATGCAGAGGCTGACATAGTGAACCTCCAAGGAGTGACAGGAGGCAACACATACT 1984

Qy      2080 CTGTGCCTGCACTGCCCCCAGGGGCAGTC---GGGGATGGGCCCCCCAGAGTGGATTTCC 2136
Db      1985 CAGTGCCTGCCGTCACCATGGACCTGCTCTCAGGAAAAGATGTGGCTGTGGAGGAGTTCC 2044

Qy      2137 CTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGGGCCAGTTTGGGGAGGTGCACC 2196
Db      2045 CCAGGAAACTCCTAACTTTCAAAGAGAAGCTGGGAGAAGGACAGTTTGGGGAGGTTCATC 2104

Qy      2197 TGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTGATTTCCCCCTTAATGTGCGTA 2256
Db      2105 TCTGTGAAGTGGAGGGAATGGAAAAATTCAAAGACAAAGATTTTGCCCTAGATGTCAGTG 2164
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Qy       2257 AGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGCCAGATGCCACCAAGAATGCCA 2316
Db       2165 CCAACCAGCCTGTCCTGGTGGCTGTGAAAATGCTCCGAGCAGATGCCAACAAGAATG--- 2221

Qy       2317 GCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTCA 2376
Db       2222 --------------CCAGGAATGATTTTCTTAAGGAGATAAAGATCATGTCTCGGCTCA 2266

Qy       2377 AGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGA 2436
Db       2267 AGGACCCAAACATCATCCATCTATTATCTGTGTGTATCACTGATGACCCTCTCTGTATGA 2326

Qy       2437 TTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAGG 2496
Db       2327 TCACTGAATACATGGAGAATGGAGATCTCAATCAGTTTCTTTCCCGCCACGAGCCCCCTA 2386

Qy       2497 ACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCT 2556
Db       2387 ATTCTTCCTCCAGCG--------------------------ATGTACGCACTGTCAGTT 2419

Qy       2557 ACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACAC 2616
Db       2420 ACACCAATCTGAAGTTTATGGCTACCCAAATTGCCTCTGGCATGAAGTACCTTTCCTCTC 2479

Qy       2617 TCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCA 2676
Db       2480 TTAATTTTGTTCACCGAGATCTGGCCACACGAAACTGTTTAGTGGGTAAGAACTACACAA 2539

Qy       2677 TCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTGC 2736
Db       2540 TCAAGATAGCTGACTTTGGAATGAGCAGGAACCTGTACAGTGGTGACTATTACCGGATCC 2599

Qy       2737 AGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAGT 2796
Db       2600 AGGGCCGGGCAGTGCTCCCTATCCGCTGGATGTCTTGGGAGAGTATCTTGCTGGGCAAGT 2659

Qy       2797 TCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCCTGTGGGAGGTGCTGATGCTCT 2856
Db       2660 TCACTACAGCAAGTGATGTGTGGGCCTTTGGGGTTACTTTGTGGGAGACTTTCACCTTTT 2719

Qy       2857 GTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGAGAACGCGGGGGAGT 2916
Db       2720 GTCAAGAACAGCCCTATTCCCAGCTGTCAGATGAACAGGTTATTGAGAATACTGGAGAGT 2779

Qy       2917 TCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGCCGCCTGCCTGCCCGCAGGGCC 2976
Db       2780 TCTTCCGAGACCAAGGGAGGCAGACTTACCTCCCTCAACCAGCCATTTGTCCTGACTCTG 2839

Qy       2977 TATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTGAGCAGCGACCACCCTTTTCCC 3036
Db       2840 TGTATAAGCTGATGCTCAGCTGCTGGAGAAGAGATACGAAGAACCGTCCCTCATTCCAAG 2899

Qy       3037 AGCTGCATCGGTTCCT 3052
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RESULT 11
US-08-336-343A-5/c
; Sequence 5, Application US/08336343A
; Patent No. 5677144
;  GENERAL INFORMATION:
;    APPLICANT:  Ullrich, Axel
;    APPLICANT:  Alves, Frauke
;    TITLE OF INVENTION:  CCK-2, A No. 5677144el Receptor Tyrosine Kinase
;    NUMBER OF SEQUENCES:  43
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Pennie & Edmonds
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/336,343A
;      FILING DATE:  08-NOV-1994
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Coruzzi, Laura A.
;      REGISTRATION NUMBER:  30,742
;      REFERENCE/DOCKET NUMBER:  7683-065
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 790-9090
;      TELEFAX:  (212) 869-9741/8864
;      TELEX:  66141 PENNIE
;  INFORMATION FOR SEQ ID NO:  5:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3157 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
US-08-336-343A-5

  Query Match             16.2%;  Score 642;  DB 1;  Length 3157;
  Best Local Similarity   56.5%;  Pred. No. 7.2e-142;
  Matches 1534;  Conservative    0;  Mismatches  975;  Indels  207;  Gaps   10;

Qy        349 TGCTGCTGCTCTTGGTGGCAAGTGGAGATGCTGACATGAAGGGACATTTTGATCCTGCCA 408
Db       2763 TGGTGCTGTTCCTGCTGCTGCCTATCTTGAGTTCTGCAAAAGCTCAGGTTAATCCAGCTA 2704

Qy        409 AGTGCCGCTATGCCCTGGGCATGCAGGACCGGACCATCCCAGACAGTGACATCTCTGCTT 468
Db       2703 TATGCCGCTATCCTCTGGGCATGTCAGGAGGCCAGATTCCAGATGAGGACATCACAGCTT 2644

Qy        469 CCAGCTCCTGGTCAGATTCCACTGCCGCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGG 528
Db       2643 CCAGTCAGTGGTCAGAGTCCACAGCTGCCAAATATGGAAGGCTGGACTCAGAAGAAGGGG 2584

Qy        529 ATGGGGCCTGGTGCCCCGCAGGGTCGGTGTTTCCCAAGGA---GGAGGAGTACTTGCAGG 585
Db       2583 ATGGAGCCTGGTGCCCTGAGATTCCAGTGGAACCTGATGACCTGAAGGAGTTTCTGCAGA 2524

Qy        586 TGGATCTACAACGACTCCACCTGGTGGCTCTGGTGGGCACCCAGGGACGGCATGCCGGGG 645
Db       2523 TTGACTTGCACACCCTCCATTTTATCACTCTGGTGGGGACCCAGGGGCGCCATGCAGGAG 2464

Qy        646 GCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGCT 705
Db       2463 GTCATGGCATCGAGTTTGCCCCCATGTACAAGATCAATTACAGTCGGGATGGCACTCGCT 2404

Qy        706 GGATGGGCTGGAAGGACCGCTGGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCTGAGG 765
Db       2403 GGATCTCTTGGCGGAACCGTCATGGGAAACAGGTGCTGGATGGAAATAGTAACCCCTATG 2344

Qy        766 GAGTGGTGCTGAAGGACCTTGGGCCCCCCATGGTTGCCCGACTGGTTCGCTTCTACCCCC 825
Db       2343 ACATTTTCCTAAAGGACTTGGAGCCGCCCATTGTAGCCAGATTTGTCCGGTTCATTCCAG 2284

Qy        826 GGGCTGACCGGGTCATGAGTGTCTGTCTGCGGGTAGAGCTCTATGGCTGCCTCTGGAGGG 885
Db       2283 TCACCGACCACTCCATGAATGTGTGTATGAGAGTGGAGCTTTACGGCTGTGTCTGGCTAG 2224

Qy        886 ATGGACTCCTGTCTTACACCGCCCCTGTGGGGCAGACAATGTATTTATCTGAGG------ 939
Db       2223 ATGGCTTGGTGTCTTACAATGCTCCAGCTGGGCAGCAGTTTGTACTCCCTGGAGGTTCCA 2164
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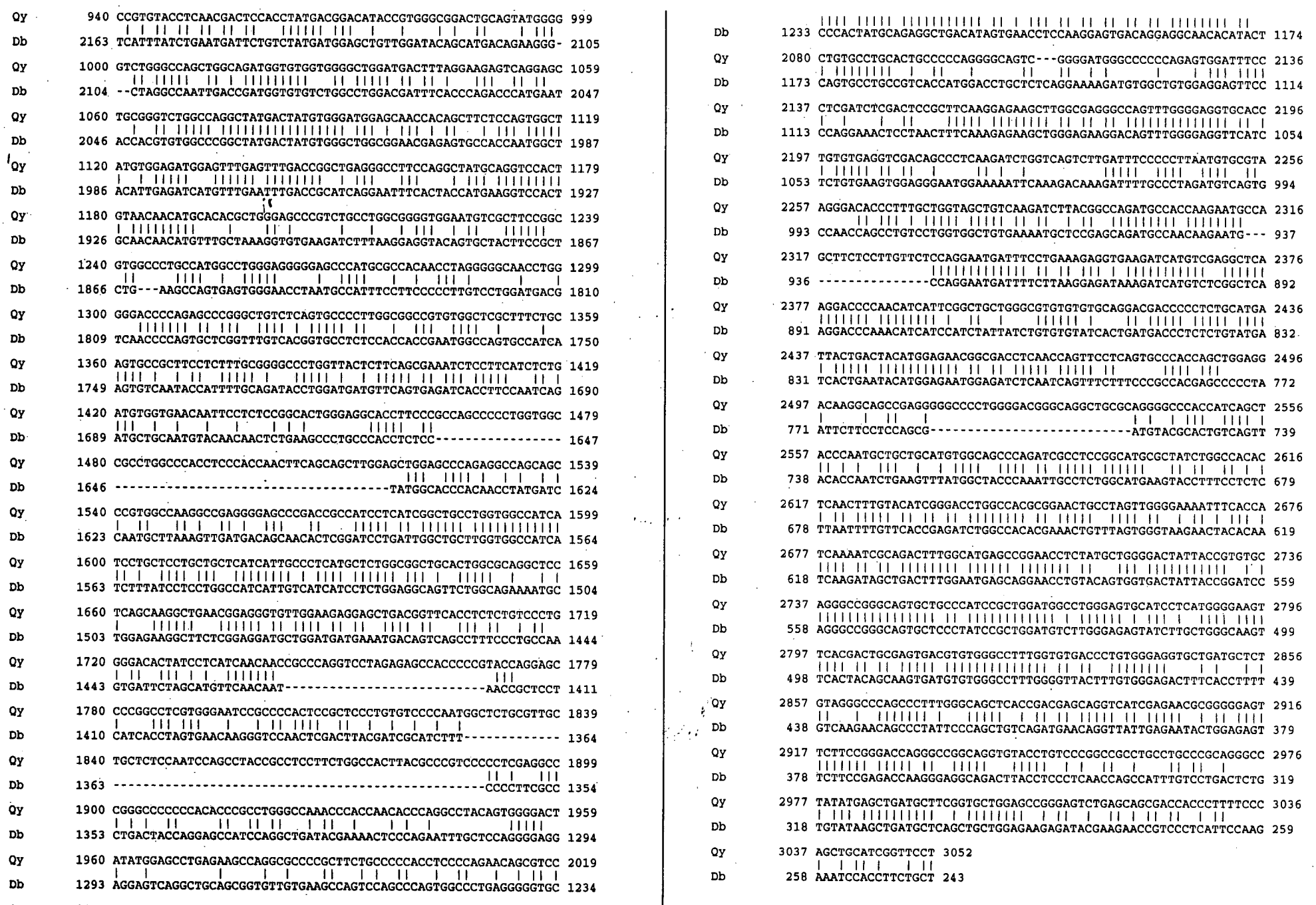

RESULT 12

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US-08-456-647B-19
; Sequence 19, Application US/08456647B
; Patent No. 5811516
;  GENERAL INFORMATION:
;    APPLICANT:  Lemke Ph.D. et al., Greg E.
;    TITLE OF INVENTION:  PROTEIN-TYROSINE KINASE GENES
;    NUMBER OF SEQUENCES:  54
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Fish & Richardson P.C.
;      STREET:  4225 Executive Square, Suite 1400
;      CITY:  La Jolla
;      STATE:  CA
;      COUNTRY:  US
;      ZIP:  92037
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/456,647B
;      FILING DATE:  02-JUN-1995
;      CLASSIFICATION:  530
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/237,401
;      FILING DATE:  02-MAY-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/884,486
;      FILING DATE:  15-MAY-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Wetherell Ph.D., John R.
;      REGISTRATION NUMBER:  31,678
;      REFERENCE/DOCKET NUMBER:  07251/007002
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (619) 678-5070
;      TELEFAX:  (619) 678-5099
;  INFORMATION FOR SEQ ID NO:  19:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3120 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
;    IMMEDIATE SOURCE:
;      CLONE:  Tyro-10
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  485..3047
US-08-456-647B-19

  Query Match           16.1%;  Score 639.8;  DB 1;  Length 3120;
  Best Local Similarity  56.5%;  Pred. No. 2.4e-141;
  Matches 1552;  Conservative    0;  Mismatches  987;  Indels  210;  Gaps   11;

Qy        348 CTGCTGCTGCTCTTGGTGGCAAGTGGAGATGCTGACATGAAGGGACATTTTGATCCTGCC 407
Db        509 CTGGTGCTGCTCCTGCTCTTGCTCATCCTGGGTTCTGCAAAAGCTCAGGTTAATCCAGCC 568

Qy        408 AAGTGCCGCTATGCCCTGGGCATGCAGGACCGGACCATCCCAGACAGTGACATCTCTGCT 467
Db        569 ATATGCCGCTATCCTCTGGGCATGTCAGGAGGCCACATTCCAGATGAGGACATCACAGCC 628

Qy        468 TCCAGCTCCTGGTCAGATTCCACTGCCGCCCGCCACAGCAGGTTGGAGAGCAGTGACGGG 527
Db        629 TCAAGTCAGTGGTCAGAATCCACGGCTGCCAAATATGGGAGGCTGGACTCTGAAGAAGGA 688

Qy        528 GATGGGGCCTGGTGCCCCGCAGGGTCGGTGTTTCCCAAGGA---GGAGGAGTACTTGCAG 584
Db        689 GATGGAGCCTGGTGTCCTGAGATTCCAGTGCAACCCGATGACCTGAAGGAATTTCTGCAG 748

Qy        585 GTGGATCTACAACGACTCCACCTGGTGGCTCTGGTGGGCACCCAGGGACGGCATGCCGGG 644
Db        749 ATTGACTTGCGAACCCTACACTTTATCACTCTTGTGGGGACCCAGGGGCGCCATGCAGGG 808

Qy        645 GGCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGC 704
Db        809 GGTCATGGCATTGAATTTGCACCCATGTACAAGATCAACTACAGTCGGGATGGCAGTCGC 868

Qy        705 TGGATGGGCTGGAAGGACCGCTGGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCTGAG 764
Db        869 TGGATCTCCTGGCGTAACCGGCATGGGAAGCAGGTGCTTGATGGAAACAGTAACCCTTAT 928

Qy        765 GGAGTGGTGCTGAAGGACCTTGGGCCCCCCATGGTTGCCCGACTGGTTCGCTTCTACCCC 824
Db        929 GATGTATTCCTGAAGGACTTGGAGCCACCCATCGTCGCCAGATTTGTTCGCCTTATCCCA 988

Qy        825 CGGGCTGACCGGGTCATGAGTGTCTGTCTGCGGGTAGAGCTCTATGGCTGCCTCTGGAGG 884
Db        989 GTCACTGACCACTCCATGAACGTGTGCATGAGGGTTGAGCTTTATGGTTGTGTCTGGCTA 1048

Qy        885 GATGGACTCCTGTCTTACACCGCCCCTGTGGGGCAGACAATGTATTTATCTGAGG----- 939
Db       1049 GATGGCTTGGTATCCTACAATGCTCCAGCTGGACAGCAGTTTGTACTCCCTGGAGGCTCC 1108

Qy        940 -CCGTGTACCTCAACGACTCCACCTATGACGGACATACCGTGGGCGGACTGCAGTATGGG 998
Db       1109 ATCATTTATCTGAATGATTCTGTCTATGATGGAGCTGTTGGGTACAGCATGACTGAAGGG 1168

Qy        999 GGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGGATGACTTTAGGAAGAGTCAGGAG 1058
Db       1169 ---CTAGGCCAGTTGACTGATGGAGTATCCGGCCTGGATGATTTTACCCAGACCCATGAA 1225

Qy       1059 CTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGAGCAACCACAGCTTCTCCAGTGGC 1118
Db       1226 TACCACGTGTGGCCTGGCTATGACTACGTGGGATGGCGGAACGAAAGTGCTACCAACGGT 1285

Qy       1119 TATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCAC 1178
Db       1286 TTCATTGAGATCATGTTTGAATTTGACCGAATCAGGAATTTTACTACCATGAAGGTCCAC 1345

Qy       1179 TGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTGGCGGGGTGGAATGTCGCTTCCGG 1238
Db       1346 TGCAACAACATGTTTGCTAAAGGTGTGAAGATTTTTAAGGAGGTCCAGTGCTACTTTCGC 1405

Qy       1239 CGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTG 1298
Db       1406 TCGG---AAGCCAGCGAGTGGGAACCCACTGCTGTCTACTTTCCCCTGGTCCTGGACGAT 1462

Qy       1299 GGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTG 1358
Db       1463 GTGAACCCCAGTGCCCGGTTTGTCACGGTGCCCCTCCACCACCGAATGGCCAGTGCCATC 1522

Qy       1359 CAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCTTCAGCGAAATCTCCTTCATCTCT 1418
Db       1523 AAGTGCCAATACCATTTTGCCGACACGTGGATGATGTTCAGCGAGATCACTTTCCAATCA 1582

Qy       1419 GATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCACCTTCCCGCCAGCCCCCTGGTGG 1478
Db       1583 GATGCTGCAATGTATAACAACTCTGGAGCCCTTCCCACCTCTCC---------------- 1626

Qy       1479 CCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGGAGCTGGAGCCCAGAGGCCAGCAG 1538
Db       1627 ------------------------------------TATGGCACCCACCACCTATGAT 1648

Qy       1539 CCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCCTCATCGGCTGCCTGGTGGCCATC 1598
Db       1649 CCCATGCTTAAAGTTGATGATAGCAACACTCGGATCCTGATTGGTTGCTTGGTGGCCATC 1708

Qy       1599 ATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCTGGCGGCTGCACTGGCGCAGGCTC 1658
Db       1709 ATCTTCATCCTGCTGGCTATCATCGTCATCATCCTGTGGAGGCAGTTCTGGCAGAAGATG 1768

Qy       1659 CTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGCTGACGGTTCACCTCTCTGTCCCT 1718
Db       1769 CTAGAAAAGGCTTCACGGAGGATGCTGGATGATGAAATGACAGTCAGCCTTTCCCTGCCC 1828
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Qy      1719 GGGGACACTATCCTCATCAACAACCGCCCAGGTCCTAGAGAGCCACCCCCGTACCAGGAG 1778
Db      1829 AGCGAGTCCAGCATGTTCAATAACAACCGCTCCTC------ATCACCAAGTGAACAGGAG 1882

Qy      1779 CCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCTGTGTCCCCAATGGCTCTGCGTTG 1838
Db      1883 TCCAACTCTACTTATGATCGAATCTTCCCCCTTCGCCCTGACTACCAGGAGCCATCCAGA 1942

Qy      1839 CTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCACTTACGCCCGTCCCCCTCGAGGC 1898
Db      1943 CTGATC---------------------------------------------------- 1948

Qy      1899 CCGGGCCCCCCCCACACCCGCCTGGGCCAAACCCACCAACACCCAGGCCTACAGTGGGGAC 1958
Db      1949 ------------------------CGAAAGCTTCCAGAGTTTGCTCCAGGAGAGGAGGA 1983

Qy      1959 TATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGCCCCCACCTCCCCAGAACAGCGTC 2018
Db      1984 GTCAGGGTGCAGTGGTGTTGTGAAGCCGGCC-----CAGCCCAATGGACCTGAGGGCGTG 2038

Qy      2019 CCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGGGCGTCACCGGGGGCAACACCTAT 2078
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Qy      2436 ATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAG 2495
Db      2441 ATCACGGAATACATGGAGAATGGAGATCTTAATCAGTTTCTTTCTCGCCACGAGCCTCTG 2500

Qy      2496 GACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGC 2555
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Qy      2556 TACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACA 2615
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Db      2591 CTCAACTTTGTCCACCGAGATCTGGCCACACGAAACTGTTTAGTGGGCAAGAATTACACC 2650

Qy      2676 ATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTG 2735
Db      2651 ATCAAGATAGCTGATTTTGGCATGAGCAGAAACCTGTACAGTGGTGATTACTACCGGATC 2710

Qy      2736 CAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAG 2795
Db      2711 CAGGGCCGGGCGGTGCTCCCCATTCGCTGGATGTCCTGGGAAAGCATCTTGCTGGGCAAA 2770

Qy      2796 TTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCCTGTGGGAGGTGCTGATGCTC 2855

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Qy      2856 TGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGAGAACGCGGGGGAG 2915
Db      2831 TGCCAGGAGCAGCCCTATTCCCAGCTGTCGGATGAGCAGGTTATCGAGAACACTGGAGAG 2890

Qy      2916 TTCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGCCGCCTGCCTGCCCGCAGGGC 2975
Db      2891 TTCTTCCGAGACCAAGGGAGGCAGATCTATCTCCCTCAACCAGCCCTTTGCCCCGACTCT 2950

Qy      2976 CTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTGAGCAGCGACCACCCTTTTCC 3035
Db      2951 GTGTATAAGCTGATGCTCAGCTGCTGGAGAAGAGAAACCAAGCACCGGCCATCCTTCCAG 3010

Qy      3036 CAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGGTGTGAATCA 3084
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RESULT 13
US-08-237-401A-19
; Sequence 19, Application US/08237401A
; Patent No. 5837448
; GENERAL INFORMATION:
; APPLICANT: Lemke Ph.D. et al., Greg E.
; TITLE OF INVENTION: PROTEIN-TYROSINE KINASE GENES
; NUMBER OF SEQUENCES: 54
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 4225 Executive Square, Suite 1400
; CITY: La Jolla
; STATE: CA
; COUNTRY: US
; ZIP: 92037
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/237,401A
; FILING DATE: 02-MAY-1994
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/884,486
; FILING DATE: 15-MAY-1992
; ATTORNEY/AGENT INFORMATION:
; NAME: Haile Ph.D., Lisa A.
; REGISTRATION NUMBER: 38,347
; REFERENCE/DOCKET NUMBER: 07251/007001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (619) 678-5070
; TELEFAX: (619) 678-5099
; INFORMATION FOR SEQ ID NO: 19:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3120 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA
; IMMEDIATE SOURCE:
; CLONE: Tyro-10
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 485..3047
US-08-237-401A-19

Query Match 16.1%; Score 639.8; DB 2; Length 3120;
Best Local Similarity 56.5%; Pred. No. 2.4e-141;
Matches 1552; Conservative 0; Mismatches 987; Indels 210; Gaps 11;

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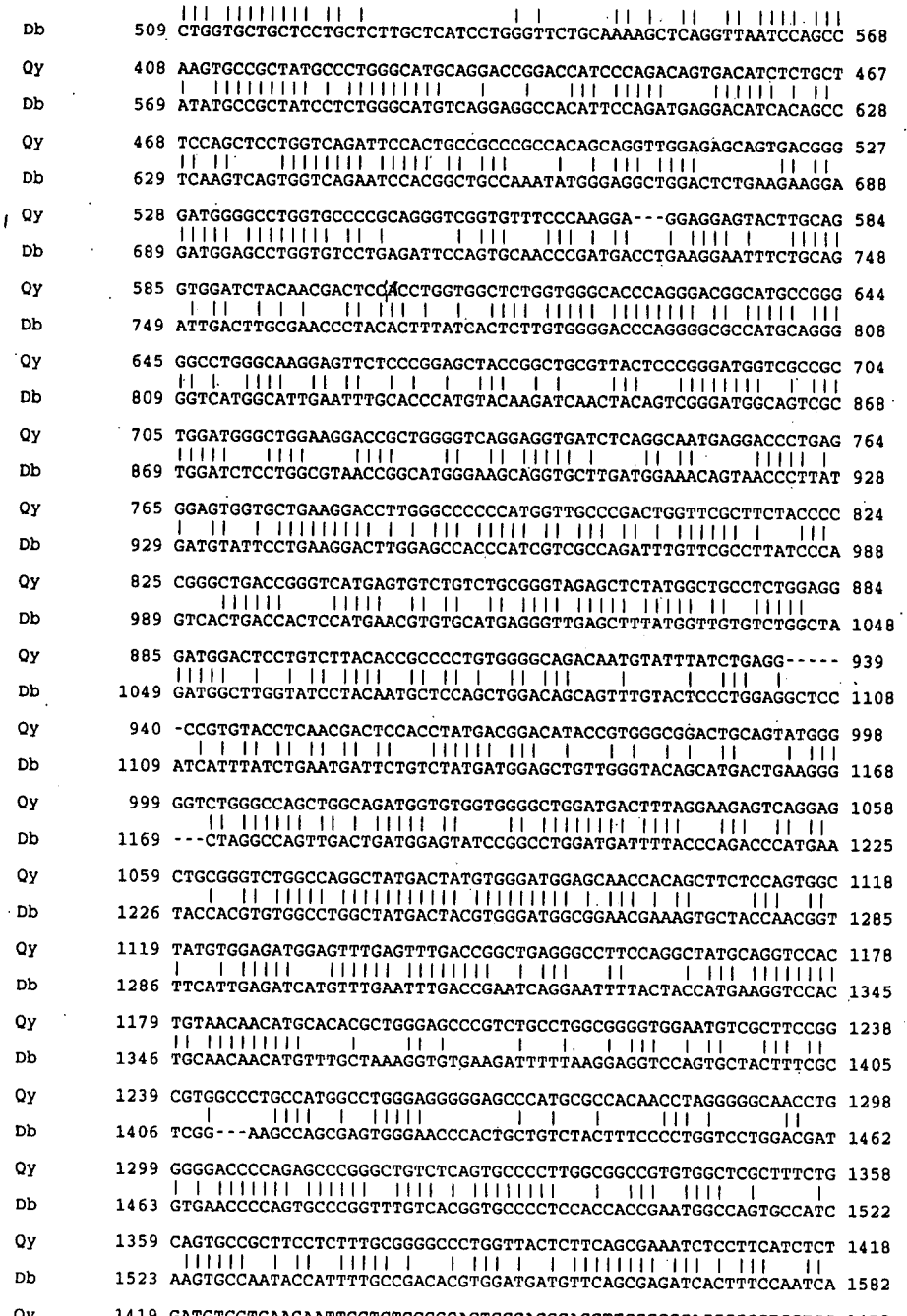
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Qy 1119 TATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCAC 1178
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Db 1346 TGCAACAACATGTTTGCTAAAGGTGTGAAGATTTTTAAGGAGGTCCAGTGCTACTTTCGC 1405
Qy 1239 CGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTG 1298
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Qy 1299 GGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTG 1358
Db 1463 GTGAACCCCAGTGCCCGGTTTGTCACGGTGCCCCTCCACCACCGAATGGCCAGTGCCATC 1522
Qy 1359 CAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCTTCAGCGAAATCTCCTTCATCTCT 1418
Db 1523 AAGTGCCAATACCATTTTGCCGACACGTGGATGATGTTCAGCGAGATCACTTTCCAATCA 1582
Qy 1419 GATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCACCTTCCCGCCAGCCCCCTGGTGG 1478
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Qy 2316 AGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTC 2375
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Db 2501 A----------GTTCCTGTTCTAGTGATGCCACAGTCAGT 2530

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Qy       2556 TACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACA 2615
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Qy       2616 CTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACC 2675
Db       2591 CTCAACTTTGTCCACCGAGATCTGGCCACACGAAACTGTTTAGTGGGCAAGAATTACACC 2650

Qy       2676 ATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTG 2735
Db       2651 ATCAAGATAGCTGATTTTGGCATGAGCAGAAACCTGTACAGTGGTGATTACTACCGGATC 2710

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; Sequence 4, Application US/08286305A
; Patent No. 5766863
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Sadick, Michael D.
;    APPLICANT:  Shelton, David L.
;    APPLICANT:  Wong, Wai Lee Tan
;    TITLE OF INVENTION:  KINASE RECEPTOR ACTIVATION ASSAY
;    NUMBER OF SEQUENCES:  11
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/286,305A
;      FILING DATE:  05-AUG-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Lee, Wendy M.
;      REGISTRATION NUMBER:  00,000
;      REFERENCE/DOCKET NUMBER:  854C1P1
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1994
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2820 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
US-08-286-305A-4

  Query Match             4.6%;  Score 182.2;  DB 1;  Length 2820;
  Best Local Similarity   59.9%;  Pred. No. 1.5e-33;
  Matches  348;  Conservative    0;  Mismatches  218;  Indels   15;  Gaps    2;

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Qy       2382 CCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGATTACT 2441
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Qy       2862 GCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGA 2902
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RESULT 15
US-08-441-104A-4
; Sequence 4, Application US/08441104A
; Patent No. 5891650
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Sadick, Michael D.
;    APPLICANT:  Shelton, David L.
;    APPLICANT:  Wong, Wai Lee Tan
;    TITLE OF INVENTION:  KINASE RECEPTOR ACTIVATION ASSAY
;    NUMBER OF SEQUENCES:  11
;    CORRESPONDENCE ADDRESS:
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;       ADDRESSEE:  Genentech, Inc.
;       STREET:  460 Point San Bruno Blvd
;       CITY:  South San Francisco
;       STATE:  California
;       COUNTRY:  USA
;       ZIP:  94080
;     COMPUTER READABLE FORM:
;       MEDIUM TYPE:  3.5 inch, 720 Kb floppy disk
;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  patin (Genentech)
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/441,104A
;       FILING DATE:  15-MAY-1995
;       CLASSIFICATION:  435
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/286305
;       FILING DATE:  05-AUG-1994
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/170558
;       FILING DATE:  20-DEC-1993
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/157563
;       FILING DATE:  23-NOV-1993
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Lee, Wendy M.
;       REGISTRATION NUMBER:  00,000
;       REFERENCE/DOCKET NUMBER:  854C1P1C2
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  415/225-1994
;       TELEFAX:  415/952-9881
;       TELEX:  910/371-7168
;   INFORMATION FOR SEQ ID NO:  4:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  2820 bases
;       TYPE:  nucleic acid
;       STRANDEDNESS:  double
;       TOPOLOGY:  linear
US-08-441-104A-4

  Query Match             4.6%;  Score 182.2;  DB 2;  Length 2820;
  Best Local Similarity   59.9%;  Pred. No. 1.5e-33;
  Matches  348;  Conservative    0;  Mismatches  218;  Indels   15;  Gaps    2;

Qy       2322 TCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTCAAGGAC 2381
Db       1994 TCCGAGAGTGCTCGGCAGGACTTCCAACGTGAGGCTGAGCTGCTCACCATGCTGCAGCAC 2053

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Qy       2502 GCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCTACCCA 2561
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Qy       2622 TTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCATCAAA 2681
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Qy       2682 ATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTGCAGGGC 2741
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Qy       2742 CGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAGTTCACG 2801

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Qy       2862 GCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGA 2902
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Search completed: May 30, 2003, 13:08:14
Job time : 176 secs
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